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UTILITY PATENT APPLICATION TRANSMITTAL (Large Entity)

(Only for new nonprovisional applications under 37 CFR 1.53(b))

Docket No. 1030-R-00

Total Pages in this Submission 171

TO THE ASSISTANT COMMISSIONER FOR PATENTS

Box Patent Application Washington, D.C. 20231

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	Application Elements	
1. ⊠ Fi	ling fee as calculated and transmitted as described below	
2. 🗵 S	pecification having pages and including the following:	
a. 🗵	Descriptive Title of the Invention	
b. 🗵	Cross References to Related Applications (if applicable)	
с. 🗌	Statement Regarding Federally-sponsored Research/Development (if applicable	э)
d. 🗆	Reference to Microfiche Appendix (if applicable)	
e. 🗵	Background of the Invention	
f. 🗵	Brief Summary of the Invention	
g. 🛚	Brief Description of the Drawings (if drawings filed)	
h. 🗵	Detailed Description	
i. 🛚	Claim(s) as Classified Below	
j. 🛚	Abstract of the Disclosure	

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	_	Application Elements (Continued)
3.	X	Drawing(s) (when necessary as prescribed by 35 USC 113)
	a.	☐ Formal Number of Sheets
	b.	
4.	X	Oath or Declaration
	a.	☐ Newly executed (original or copy) ☐ Unexecuted
	b.	☐ Copy from a prior application (37 CFR 1.63(d)) (for continuation/divisional application only)
	C.	☑ With Power of Attorney ☐ Without Power of Attorney
	d.	☐ <u>DELETION OF INVENTOR(S)</u> Signed statement attached deleting inventor(s) named in the prior application, see 37 C.F.R. 1.63(d)(2) and 1.33(b).
5.		Incorporation By Reference (usable if Box 4b is checked) The entire disclosure of the prior application, from which a copy of the oath or declaration is supplied under Box 4b, is considered as being part of the disclosure of the accompanying application and is hereby incorporated by reference therein.
6.		Computer Program in Microfiche (Appendix)
7.	\boxtimes	Nucleotide and/or Amino Acid Sequence Submission (if applicable, all must be included)
	a.	☑ Paper Copy
	b.	☑ Computer Readable Copy (identical to computer copy)
	C.	☑ Statement Verifying Identical Paper and Computer Readable Copy
		Accompanying Application Parts
8.		Assignment Papers (cover sheet & document(s))
9.		37 CFR 3.73(B) Statement (when there is an assignee)
10.		English Translation Document (if applicable)
11.	X	Information Disclosure Statement/PTO-1449 Copies of IDS Citations
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Docket No. 1030-R-00

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APPLICATION FOR UNITED STATES LETTERS PATENT

FOR

A METHOD FOR THE IDENTIFICATION OF ANESTHETICS

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BACKGROUND OF THE INVENTION

This patent application is a continuation-in-part of pending application Serial No. 09/144,914 filed September 1, 1998, which is incorporated by reference in its entirety. This patent application also claims the priority of U.S. Provisional Application No. 60/119,727, filed February 12, 1999.

1. Field of the Invention

The invention relates to a method for identifying substances that are capable of acting as anesthetics.

2. Background of the Related Art

Volatile anesthetics are a remarkable class of agents producing a safe, reversible state of unconsciousness with concurrent amnesia and analgesia. They have hyperpolarizing action on mammalian neurons. They activate an inhibitory synaptic K^+ current $(I_{K(An)})$ in molluscan pacemaker neurons which has been proposed to have an important role in general anaesthesia.

Volatile anesthetics hyperpolarize frog motor neurons, rat hippocampal neurons, guinea pig thalmic neurons and human cerebral cortex neurons. Therefore, it has been proposed that the molecular mechanism of volatile anesthetics involves an action on a specific class of K+ channels. The fact that a particular inhibitory synaptic K+ current, $I_{K(An)}$, reversibly activated by volatile agents is present in anesthetic-sensitive molluscan pacemaker neurons, but absent in insensitive neurons has made it a very attractive candidate as a target for these important pharmacological agents. $I_{K(An)}$ behaves as a background channel; it is not voltage-gated, it activates immediately, and it does not inactivate with time. $I_{K(An)}$ obeys the Goldman-Hodgkin-Katz constant field equation and is resistant to the classical K+ channel blockers, tetraethylammonium and 4-aminopyridine.

We recently identified a novel family of mammalian K+ channels with a unique structural

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motif consisting of two pore domains in tandem and four transmembrane segments. The four members in this family have been classified as TWIK-1, TASK, TREK-1, and TRAAK, which are shown in Fig. 1A. TWIK-1 has been previously shown to dimerize, implying that a functional channel is formed by at least two subunits. Heteromultimerization does not occur between the four members of this novel family as tested in Sf9 cells expressing various combinations of these channels (unpublished data). Three members of this family, mouse TREK-1, mouse TRAAK, and human TASK, encode for background outward-going K+rectifiers with properties resembling those of $I_{K(An)}$. TRAAK and TREK-1 are directly opened by arachidonic acid and other polyunsaturated fatty acids, while TASK encodes a resting K+ channel which is controlled by external pH variations near physiological pH. TREK-1 and TASK are expressed in many tissues and are particularly abundant in the brain and in the heart, whereas TRAAK is selectively expressed in the central nervous system. Neuronal expression of these channels is detected at high levels in the cortex, cerebellum, hippocampus and olfactory bulb, and cardiac expression is detected in both the myocardium and connective tissues.

BRIEF DESCRIPTION OF THE FIGURES

Fig. 1 shows the results of experiments with chloroform demonstrating that chloroform selectively activates TREK-1: A, a schematic showing the two pore domains and four transmembrane domains of the 2P domain potassium K+ channels, TREK-1 TRAAK, TWIK-1 and TASK. The four transmembrane domain segments are indicated by TM1 through TM4, and the two pore regions are indicated by P1 and P2. The phylogenetic tree indicates the three subfamilies. TWIK-1 is an inward rectifier K+ channel and TASK is a background rectifier K+ channel inhibited

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by external acidosis. Both TREK-1 and TRAAK are background outward rectifier K+ channels opened by arachidonic acid; B, whole cell patch clamp experiments showing that 0.8 mM chloroform strongly and reversibly activates TREK-1 expressed in transfected COS cells. The mock condition is the wild-type (empty) expression vector. The effects of chloroform were investigated on K+ currents elicited in the whole cell configuration during a voltage ramp of one second in duration from a holding potential of -80 mV as illustrated in the inset for TREK-1; C, 0.8 mM chloroform induced a typical background current characterized by an outward-going rectification which reverses at the predicted value for E_{K+}; Chloroform activated TREK-1 currents in physiological and symmetrical K+ gradients were examined. Voltage ramps of one second in duration from a holding potential of -80 mV in both K+ conditions and in the presence of 0.8 mM chloroform were digitally subtracted from ramps in control conditions; D, 0.8 mM chloroform reversibly and reproducibly hyperpolarized COS cells expressing TREK-1; E, dose-dependence of TREK-1 activation. The inset illustrates the effect of 0.8 mM chloroform on TREK-1 current measured at a holding potential of 0 mV. The number of cells in each experimental condition is indicated.

Fig. 2 Halothane is a common activator of TREK-1 and TASK. A, comparative effects of 1 mM halothane on the 2P domain K+ channel activities. The mock condition is the wild-type (empty) expression vector; B, halothane (1 mM) stimulates TASK channel activity elicited in the whole cell configuration during voltage ramps of one second in duration from a holding potential of -80 mV; C, halothane (1 mM)-activated TASK currents in physiological and symmetrical K+ gradients. Voltage ramps of one second in duration from a holding potential of -80 mV in both K+ conditions and in the presence of 1 mM halothane are digitally subtracted from ramps in control

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conditions; D, dose-effect curve of halothane on TREK-1 channel activation; F, halothane (1 mM) reversibly activates TASK at a holding potential of 0 mV. The number of cells in each experimental condition is indicated.

Fig. 3 Isoflurane and diethyl ether differentially activate TREK-1 and TASK. A, comparative effects of 2 mM isoflurane (A) and 0.6 mM diethyl ether (B) on the 2P domain K+ channels. The mock condition is the wild-type (empty) expression vector. The effects of the anesthetics are investigated on K+ currents elicited in whole cell configuration during voltage ramps of one second in duration from a holding potential of -80 mV as illustrated in insets for TREK-1. The number of cells in each experimental condition is indicated.

Fig. 4 Volatile anesthetics stimulate TREK-1 and TASK in the excised patch configurations. A, effects of increasing concentrations of halothane on TREK-1 channel activity in an outside-out patch. In an outside-out patch, halothane reversibly opens a 48 pS TREK-1 channel in a dose-dependent fashion. The holding potential is 0 mV and applications of halothane are indicated by horizontal bars; B, effect of 0.8 mM chloroform on the I-V curve of TREK-1 in an outside-out patch. The I-V curve is performed with a voltage ramp of one second in duration from a holding potential of -80 mV; C, kinetics of activation of TREK-1 by 0.8 mM chloroform. The I-V curve of the chloroform-sensitive current in an outside-out patch shows the characteristic outward-going rectification; D, halothane (1 mM) induces TASK channel opening in an inside-out patch. The holding potential is 0 mV and channel activities before, during, and after addition of 1 mM halothane are illustrated from left to right. Halothane reversibly opens a 12 pS TASK channel.

Fig. 5 Functional characterization of the human TREK-1 (hTREK-1). Transiently transfected COS cells expressing hTREK-1 are voltage-clamped using the whole cell patch clamp

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configuration. A, The basal TREK-1 current is recorded in physiological K conditions (5 mM) and in symmetrical K conditions (155 mM). The cells are held at -80 mV and voltage ramps of 800 ms in duration are applied from -130 mV to 100 mV. B, hTREK-1 is stimulated by the addition of 10 μ M arachidonic acid in the bath. Same protocol as in A. C, hTREK-1 is opened by increasing concentrations of halothane (as indicated). D, In the inside-out patch configuration hTREK-1 is opened by a membrane stretch of -66 mmHg. The patch is voltage clamped at 0 mV.

SUMMARY OF THE INVENTION

The invention relates to human TREK-1, its nucleotide and amino acid sequence. The invention further relates to murine TREK-1, its nucleotide and amino acid sequence.

Thus, the invention relates to a method for the identification of substances that activate potassium transport through a potassium transport protein, including TREK-1 and TASK. The potassium transport proteins activated in the method exhibit outward-going rectification. Specifically, the method involves contacting a substance to be tested with a mammalian transport protein and examining the potassium transport activity of the potassium transport proteins. A positive result, activation of potassium transport, correlates with a substance which produces anesthesia.

DETAILED DESCRIPTION OF THE INVENTION

The invention is an isolated nucleic acid molecule encoding the human TREK-1 channel (SEQ ID NO:1). The invention is also embodied in the isolated human TREK-1 protein (SEQ ID NO:2). The nucleic acid and deduced amino acid sequence is shown in SEQ ID NO:1.

The invention also encompasses the isolated nucleic acid molecule encoding the murine TREK-1 channel (SEQ ID NO:3). The isolated protein, murine TREK-1 (SEQ ID NO:4), is also

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encompassed by the invention.

The method involves contacting a test substance with a potassium transport protein *in vitro* and determining whether there has been activation of potassium transport.

As used in the method, cells which express a potassium transport protein are used in the presence of the test substance. The test substance is a substance which will have certain properties when used in a mammal as a volatile inhalant. These properties may include the induction of a safe, reversible state of unconsciousness, amnesia and analgesia.

The cells expressing the potassium transport protein may transiently express the protein or constitutively express the protein. The cells may be of any type which can express the protein in appropriate conformation to allow for the transport of potassium. Examples of such cells include, mammalian cells, vertebrate cells, and invertebrate cells. Examples of mammalian cells suitable for use in the invention include, but are not limited to, cells of neuronal origin, fibroblasts, myocardial cells, COS cells, Chinese hamster ovary (CHO) cells, embryonic kidney cells, fibroblasts, HELA cells, and the like. Examples of suitable non-mammalian vertebrate cells include, but are not limited to frog oocytes, such as *Xenopus laevis* oocytes, and the like. Suitable invertebrate cells include, but are not limited to insect cells, such as *Spodoptera frugiperda* (Sf9) cells, and the like. Any cell known in the art which may be transfected to transiently or constitutively express the transport protein are suitable for use in the present invention.

The potassium transport proteins which are suitable for use in the invention include TREK-1 (SEQ ID NO:2 and SEQ ID NO:4) and TASK (SEQ ID NO:5), which may be derived from any mammalian source, such as rat, mouse, or human. The molecular sequence of TREK-1 may be the human TREK-1 as shown in SEQ ID NO:2, or an amino acid sequence that is substantially identical

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to SEQ ID NO:2. By "substantially identical" it is understood that amino acid substitutions may be made such that the overall conformation of the potassium transport protein is not significantly altered: the protein remains active as a potassium transport protein.

A suitable substantially identical protein is a protein having an amino acid sequence that is generally at least 90% identical to the amino acid sequence of human TREK-1 (SEQ ID NO:2). More preferably, the protein is at least 95% identical to SEQ ID NO:2. Most preferably, the amino acid sequence is at least 99% identical to SEQ ID NO:1.

The cells used in the method of the present invention express the potassium transport protein expressed on their surface, either constitutively or transiently. Introduction of the nucleic acid into the cells so that the protein is expressed may be by any known method such as transfection of an appropriate nucleic acid construct into the cells, microinjection of RNA encoding the protein, and the like. Many different protocols are known in the art. Two methods are briefly described herein, however, it will be appreciated by one of ordinary skill in the art that many modifications and substitutions may be made to these methods without departing from the spirit and the scope of the invention.

The coding sequence of the potassium transport protein may be inserted between the noncoding sequences 5' and 3' of a *Xenopus laevis* protein (such as globin) in an appropriate vector, such as pEXO. The construct is introduced into an appropriate cell type to replicate the vector and/or to transcribe RNA. Alternatively, the vector may be used as a template for in vitro transcription. A complementary RNA (cRNA) is transcribed and injected into a cell, such as a *Xenopus* oocyte. Such a procedure may be performed in a 0.3 ml perfusion chamber, wherein single oocytes are impaled on two standard glass microelectrodes (0.5-2.0 MW) charged with 3 M KCl and maintained under

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voltage clamp with a Dagan TEV200 amplifier, The bath solution contains 98 mM KCl,1.8 mM CaCl₂, 2 mM MgCl₂, and 5 mM HEPES at pH 7.4 with KOH.

Alternatively, functional expression of the potassium channel may be accomplished by transfection of insect cells, such as *Spodoptera frugiperda* (Sf9) cells. Briefly a suitable vector, such as pVL1392 may be used and the coding sequence for the potassium transport protein may be inserted into the vector in-frame so that expression of the potassium transport protein may be expressed. The coding sequence for the potassium transport protein may be obtained by any convenient method, such as by PCR or by digesting a plasmid containing the potassium transport protein coding sequence with appropriate restriction endonuclease(s) for subsequent ligation into the pVL1392 vector. Similarly, the amplified product of the PCR may be digested with restriction enzymes and ligated into the vector. Transfection of SF9 cells may be performed by the manufacturer's protocol (Pharmingen).

Alternatively, functional expression of the potassium channel may be by transient transfection of cells such as COS cells whereby COS cells are seeded at a density of 20,000 cells per 35 mm dish. Cells are transfected with expression vector, such as the pIRES-CD8 vector, comprising the nucleic acid molecules encoding the desired potassium channel protein. The cells may be transfected by any method known in the art, such as the DEAE dextran protocol, Ca₂PO₄ precipitation, or electroporation, for example. The transfected cells, expressing the desired potassium channel, or induced to express the desired potassium channel may then be used in the method of the invention and the cells may be assayed for the transport of potassium.

The invention will be described in greater detail with reference to the examples which are provides to illustrate the invention. The examples are not to be construed to be limiting as to the

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scope of the invention, which is set forth in the appended claims.

EXAMPLES

1. Cloning of the human TREK-1 channel

The TREK-1 channel was cloned by degenerate PCR technology using previously characterized murine sequence. Although such technology is extensively described in the literature and is familiar to those of ordinary skill in the art, briefly, degenerate oligonucleotide primers selected to amplify a region of murine TREK-1 were synthesized and placed in a polymerase chain reaction amplification using human DNA with appropriate buffer, nucleotides and DNA polymerase. The reaction is cycled through temperature stages for denaturation of DNA (generally about 94 °C), annealing of primers to DNA template (this temperature can be varied to optimize amplification and can be based on many factors, including primer length and GC content), and extension of DNA polymerization by DNA polymerase (generally at the optimum temperature for the activity of the DNA polymerase which is usually about 72 °C). The amplified DNA fragment may be isolated and cloned into a plasmid vector for subsequent sequence analysis, or the amplified DNA may be directly sequenced by known methods.

Due to the degeneracy of the DNA code, it will be well understood to one of ordinary skill in the art that substitution of nucleotides may be made without changing the amino acid sequence of the protein. Therefore, the invention includes any nucleic acid sequence for the human TREK-1 channel that encodes the amino acid sequence determined for murine TREK-1 (SEQ ID NO:2). Moreover, it is understood in the art that for a given protein's amino acid sequence, substitution of certain amino acids in the sequence can be made without significant effect on the function of the protein. Such substitutions are known in the art as "conservative substitutions." The invention

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encompasses human TREK-1 proteins that contain conservative substitutions, wherein the function of the protein is not altered. Generally, the identity of such an mutant TREK-1 will be at least 90% identical to SEQ ID NO:2. Preferably, the mutant TREK-1 will be at least 95% identical to SEQ ID NO:2. More preferably, the mutant TREK-1 will be at least 97% identical to SEQ ID NO:2. Most preferably, the mutant TREK-1 will be at least 99% identical to SEQ ID NO:2.

2. The sequencing of the murine TREK-1 channel

The sequence for murine TREK-1, which is a corrected form of murine TREK-1 reported earlier, is shown in SEQ ID NO:4. There is a longer open reading frame than originally reported, producing a protein with a deduced amino acid sequence of 411 amino acids. Due to the degeneracy of the DNA code, it will be well understood to one of ordinary skill in the art that substitution of nucleotides may be made without changing the amino acid sequence of the protein. Therefore, the invention includes any nucleic acid sequence for the murine TREK-1 channel that encodes the amino acid sequence determined for murine TREK-1 (SEQ ID NO:4). Moreover, as is the case with human TREK-1, it is understood in the art that for a given protein's amino acid sequence, substitution of certain amino acids in the sequence can be made without significant effect on the function of the The invention encompasses murine TREK-1 proteins that contain conservative protein. substitutions, wherein the function of the protein is not altered. Generally, the identity of such an mutant TREK-1 will be at least 90% identical to SEQ ID NO:4. Preferably, the mutant TREK-1 will be at least 95% identical to SEQ ID NO:4. More preferably, the mutant TREK-1 will be at least 97% identical to SEQ ID NO:4. Most preferably, the mutant TREK-1 will be at least 99% identical to SEQ ID NO:4.

3. Functional expression of the human TREK-1 channel

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The functional properties of the human TREK-1 channel were studied on the basis of transfected COS cells temporarily expressing the protein. Like the murine channel, the human channel is selective for potassium and is activated by *cis*-unsaturated fatty acids, volatile anesthetics and by stretching of the plasma membrane. The selectivity of the TREK-1 currents for potassium is shown in Fig. 5A. The currents are recordings in the whole-cell configuration of the patch-clamp technique (imposed potential gradient from -130 to +100 mV). The inverse potential of the currents follow the equilibrium potential of potassium when the extracellular concentration of potassium ions climbs from 5 to 155 mM. Fig. 5B shows that the application of arachidonic acid (10 μM) induced activation of the TREK-1 currents. Fig. 5C shows that the application of volatile anesthetics (halothane in this case) at the concentrations employed in general anesthesia induced activation of the TREK-1 channel. Fig. 5D shows that the TREK-1 currents recorded in an excised patch (insideout configuration) are mechanosensitive. When a pressure of -66 mmHg is applied on the membrane, the TREK-1 currents are activated in a reversible manner.

4. Electrophysiological recording

COS cell transfection, culture and electrophysiology are well known in the art and are described in the literature such as in the references cited.

As performed herein, COS cells were seeded at a density of 20,000 cells per 35 mm dish 24 hours prior to transfection. Cells were transfected by the DEAE dextran protocol (1 µg DNA per plate). Mouse TREK-1 (GenBank Accession No. U73488), human TASK (GenBank Accession No. AF006823) and mouse TRAAK (GenBank Accession No. AF056492) fragments were amplified by polymerase chain reaction (PCR) and subcloned into the pIRES-CD8 vector. Transfected cells were visualized 48 hours after transfection using the anti-CD8 antibody coated beads method. For whole

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cell and excised patch experiments, the internal solution was 150 mM KCl, 3 mM MgCl₂, 5mM EGTA and 10 mM HEPES at pH 7.2 with KOH and the external medium contained 150 mM NaCl, 5 mM KCL, 3 mM MgCl₂, 1 mM CaCl₂, 10 mM HEPES at pH 7.4 with NaOH. Cells were continuously superfused with a microperfusion system during the time course of the experiments (0.1 ml/min) performed at room temperature. A RK300 patch clamp amplifier (Biologic, Grenoble, France) was used for whole cell as well as single channel recordings. Ionic currents were monitored and recorded continuously using a DAT recorder (Biologic, Grenoble, France). Subsequently, data were replayed and sampled using pClamp software. Data analysis were performed using clampfit (pClamp) for whole cell recording as well as Biopatch (Biologic) software for single channel recordings. Membrane capacitance was measured during a 10 V hyperpolarizing step from a holding potential of -80 mV. Student's t test was used for statistical analysis (P<0.001).

5. Anesthetics delivery and concentration

General anesthetics were directly dissolved in saline solutions. All experiments were performed at room temperature (24°C). Solutions were prepared every 3 hours in gas-tight bottles as stock solutions (calculated concentrations of 5 10 mM). Serial dilutions were prepared just prior to the electrophysiological experiment. 2.5 ml of each experimental solution at the desired concentration was placed in a syringe connected to the experimental superfusion system. The electrophysiological measurements were performed within 45 minutes.

The actual concentrations of anesthetics were subsequently determined by means of a gas chromatography method (HP 6890 equipped with a DB624 column) using FID detection. Sampled (2.5 ml of solution) were collected prior to (t_0) and after perfusion (t_{45}) through the experimental setup. Solutions were collected using gas impermeable tubing and stored in sealed glass containers

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at 4°C for subsequent analysis. Samplings and measurements were performed in duplicate. Actual concentrations of anesthetics were determined by multiplying the calculated concentration by the ratio t_{45}/t_0 (chloroform: 0.16; halothane: 0.37; isoflurane: 0.76; and diethyl ether: 0.57). In the dose effect curves, the threshold concentrations were estimated as concentrations producing an increase higher than 10% in current amplitude.

RESULTS

The nucleic acid sequence corresponding to the open reading frame of the human TREK-1 channel is shown in SEQ ID NO:1. The sequence of 1236 nucleotides encodes a protein of 411 amino acids. The conservation between human and murine proteins is very high, exceeding 99% homology.

TASK and TREK-1, two mammalian 2P domain K+ channels which have similar properties to $I_{K(An)}$ are activated by volatile general anesthetics. Chloroform, diethyl ether, halothane and isoflurane activated TREK-1, while only halothane and isoflurane activated TASK. The C-terminal regions of both TREK-1 and TASK are critical for anesthetic activation. Thus, both TREK-1 and TASK are important target sites for these agents.

In whole patch clamp experiments shown in Fig. 1B, chloroform strongly and reversibly activated TREK-1 expressed in transfected COS cells, while it slightly depressed TASK and did not effect TRAAK. As shown in Fig. 1C, chloroform induced a typical background current, characterized by an outward-going rectification which reverses at the predicted value for E_K^+ as demonstrated by the shift of the reversal potential to 0 mV in symmetrical K^+ conditions. No current activation was observed in mock-transfected cells (Fig. 1B). Fig. 1D shows that chloroform also reversibly and reproducibly hyperpolarized COS cells expressing TREK-1 (but not mock-transfected

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cells, not shown; n=5). TREK-1 activation was dose dependent, with a threshold concentration for activation of 500 μ M, as shown in Fig. 1E. The inset of Fig. 1E shows that current activation, measured at a holding potential of 0 mV, started rapidly, but steady-state activation was barely reached after one minute in the presence of chloroform.

Both TREK and TASK, but not TRAAK, were opened by halothane. As shown in Fig. 2B and 2C, halothane-induced current displayed an outward-going rectification and revered at the predicted value for E_K^+ . The dose-effect curves (shown in Fig. 2D and Fig. 2E) demonstrated that the threshold concentrations for halothane on TREK-1 and TASK are 400 μ M and 200 μ M, respectively. The effects of halothane on TASK were rapid completely reversible. Opening of TASK was faster compared to the opening of TREK-1 by chloroform (compare Fig. 2F with Fig. 1E, inset). These results suggest that the components of the activation of both channels by anesthetics may be mediated by different molecular mechanisms.

Isoflurane, like halothane, activated both TREK-1 and TASK channels without altering TRAAK, as shown in Fig. 3A. Diethyl ether, like chloroform, opened TREK-1, but did not affect TRAAK. Diethyl ether also decreased TASK activity.

To demonstrate that activation by volatile anesthetics does not occur via second messenger pathways, experiments were conducted on excised patches. Fig. 4A shows that in an excised outside-out patch, halothane reversibly, and in a dose dependent manner, opened a 48 pS TREK-1 channel. No channel activity was observed in the absence of the anesthetic, suggesting that halothane converts inactive channels into active channels. The I-V curve of the chloroform-sensitive current in an outside-out patch shows the characteristic outward-going rectification previously observed in the macroscopic whole cell conditions, as shown in Figs. 4B-4C and Fig. 1B (inset). As

shown in Figs. 4D-4E, in the inside-out patch configuration, halothane reversibly opened a 12 pS TASK channel. In the absence of anesthetics, a single TASK channel was opened, as shown in the left panels of Figs. 4D-4E. The addition of halothane induced the opening of a second channel (see Figs. 4D-4E middle panel) which closed again after washing (see right panel of Figs. 4D and 4E). All the data, taken together, demonstrate that volatile general anesthetics open TASK and TREK-1 channels and that these effects are likely to be direct and independent of second messengers.

TREK-1 and TASK are probably critical channels for the action of general volatile anesthetics. Opening of these K+ channels along with the known modulation of neurotransmitter receptors will probably explain general anesthetic action. Effects of volatile anesthetics on ligand-gated ion channels such as GABA_A receptors are probably particularly important at the post-synaptic level. Opening of background TREK-1 and TASK channels by volatile anesthetics might be very important both at the pre-synaptic ($I_{K(An)}$) and at the post-synaptic level. An activation of only a small fraction of these $I_{K(An)}$ channels will be expected to have significant effects on membrane polarization and consequently potentially important effects in both pre- and post-synaptic functions. TREK-1 and TASK channels are expressed in neuronal cells but they are also expressed in other tissues, and particularly in the heart. It is, therefore, not surprising that volatile anesthetics have depressive side-effects on heart function. These effects include a slowing of the heart rate and negative inotropic effects and are fully compatible with an exaggerate opening of background K+ channels by volatile anesthetics. The same considerations might very well explain ventilatory depression.

What is claimed is:

- 1. An isolated nucleic acid molecule encoding the human TREK-1 potassium channel comprising the nucleic acid sequence of SEQ ID NO:1.
- 2. An isolated nucleic acid molecule encoding a human TREK-1 potassium channel, wherein said TREK-1 potassium channel comprises the amino acid sequence of SEQ ID NO:2.
- 3. An isolated human TREK-1 potassium channel comprising the amino acid sequence of SEQ ID NO:2.
- 4. The isolated human TREK-1 potassium channel of claim 3 comprising an amino acid sequence that is at least 99% identical to SEQ ID NO:2.
- 5. The isolated human TREK-1 potassium channel of claim 3 comprising an amino acid sequence that is at least 97% identical to SEQ ID NO:2.
- 6. The isolated human TREK-1 potassium channel of claim 3 comprising an amino acid sequence that is at least 95% identical to SEQ ID NO:2
- 7. An isolated nucleic acid molecule encoding the murine TREK-1 potassium channel comprising the nucleic acid sequence of SEQ ID NO:3.
- 8. An isolated nucleic acid molecule encoding a murine TREK-1 potassium channel, wherein said TREK-1 potassium channel comprises the amino acid sequence of SEQ ID NO:4.
- 9. The isolated murine TREK-1 potassium channel of claim 8 comprising the amino acid sequence of SEQ ID NO:4.
- 10. The isolated murine TREK-1 potassium channel of claim 8 comprising an amino acid sequence that is at least 99% identical to SEQ ID NO:4.

- 11. The isolated murine TREK-1 potassium channel of claim 8 comprising an amino acid sequence that is at least 97% identical to SEQ ID NO:4.
- 12. The isolated murine TREK-1 potassium channel of claim 8 comprising an amino acid sequence that is at least 95% identical to SEO ID NO:4.
- 13. A method for identifying substances having anesthetic properties, wherein said substances produce a safe, reversible state of unconsciousness with concurrent amnesia and analgesia in a mammal upon inhalation comprising:
 - (a) contacting said substance with a mammalian potassium transport protein, wherein said potassium transport protein exhibits outward-going potassium rectification; and
 - (b) determining the potassium transport activity of said potassium transport protein, wherein an activation of potassium transport is indicative of said substance having said anesthetic properties.
- 14. The method of claim 13, wherein said potassium transport protein is TASK.
- 15. The method of claim 13, wherein said potassium transport protein is TREK-1.
- 16. The method of claim 15, wherein said TREK-1 comprises the amino acid sequence selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:4.
- 17. The method of claim 15, wherein said mammalian potassium transport protein is a chimeric molecule comprising at least a portion of murine TREK-1 and at least a portion of human TREK-1 wherein said chimeric molecule comprises 2P domains and at least 4 transmembrane domains.
- 18. A method for identifying substances having anesthetic properties, wherein said substances produce a safe, reversible state of unconsciousness with concurrent amnesia and analgesia

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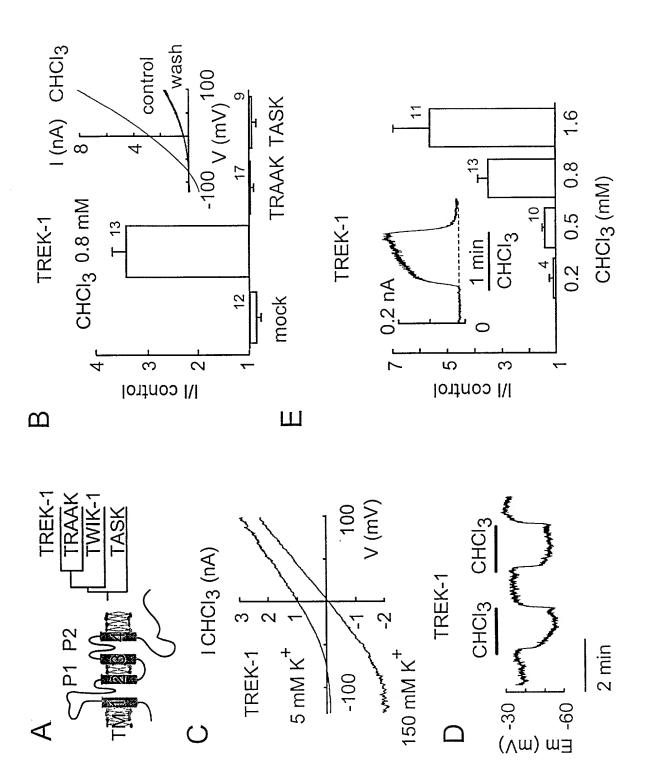
in a mammal upon inhalation comprising:

- (a) contacting said substance with COS cells, wherein said COS cells are transfected with a nucleotide vector comprising a nucleic acid molecule encoding TREK-1, wherein said COS cells transiently express said TREK-1 on a surface of said COS cells, and wherein TREK-1 exhibits outward-going potassium rectification; and
- (b) determining the potassium transport activity of said TREK-1 wherein an activation of potassium transport is indicative of said substance having said anesthetic properties.
- 19. The method of claim 18, wherein said TREK-1 comprises an amino acid sequence that is at least 95% identical to SEQ ID NO:2.
- 20. The method of claim 18, wherein said TREK-1 comprises an amino acid sequence that is at least 95% identical to SEQ ID NO: 4.
- 21. The method of claim 18, wherein said TREK-1 comprises a chimeric molecule comprising at least a portion of human TREK-1 and at least a portion of murine TREK-1.
- 22. A method for identifying substances having anesthetic properties, wherein said substances produce a safe, reversible state of unconsciousness with concurrent amnesia and analgesia in a mammal upon inhalation comprising:
 - (a) contacting said substance with transfected cells, wherein said transfected cells are transfected with a nucleotide vector comprising a nucleic acid molecule encoding TASK, wherein said transfected cells transiently express said TASK on a surface of said transfected cells, and wherein TASK exhibits outward-going potassium rectification; and
 - (b) determining the potassium transport activity of said TASK wherein an activation

- of potassium transport is indicative of said substance having said anesthetic properties.
- 23. The method of claim 22, wherein said TASK comprises an amino acid sequence that is at least 95% identical to SEQ ID NO:5.
- 24. The method of claim 22, wherein said TASK is a chimeric molecule comprising at least a portion of TASK and at least a portion of another potassium transport protein selected from the group consisting of human TREK-1 and murine TREK-1, wherein said chimeric molecule comprises 2P domains and at least 4 transmembrane domains.
- 25. The method of claim 22, wherein said transfected cells are selected from the group consisting of COS cells, HELA cells, *Spodoptera* cells, *Xenopus* oocytes, embryonic kidney cells, Chinese hamster ovary cells, and fibroblasts.

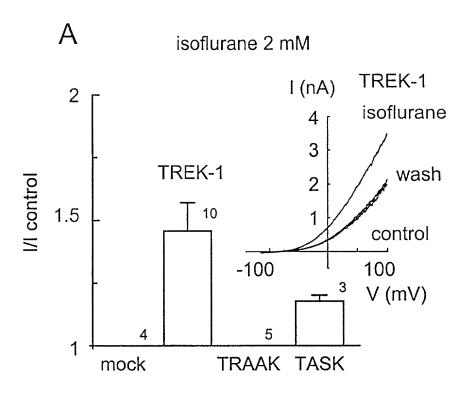
ABSTRACT OF THE DISCLOSURE

Human and mouse TREK-1 potassium transport proteins are disclosed as well a method for using potassium transport proteins for identifying substances having anesthetic properties, such as producing a safe, reversible state of unconsciousness with concurrent amnesia and analgesia in a mammal upon inhalation, in a method comprising (a) contacting the test substance with a mammalian potassium transport protein, wherein said potassium transport protein exhibits outwardgoing potassium rectification; and (b) determining the potassium transport activity of the potassium transport protein, wherein an activation of potassium transport is indicative of the test substance having anesthetic properties.



Figures 1A -1E

Figure 2A-2F



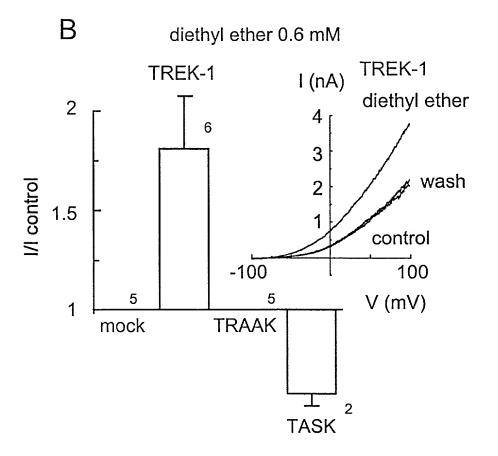


Figure 3A-B

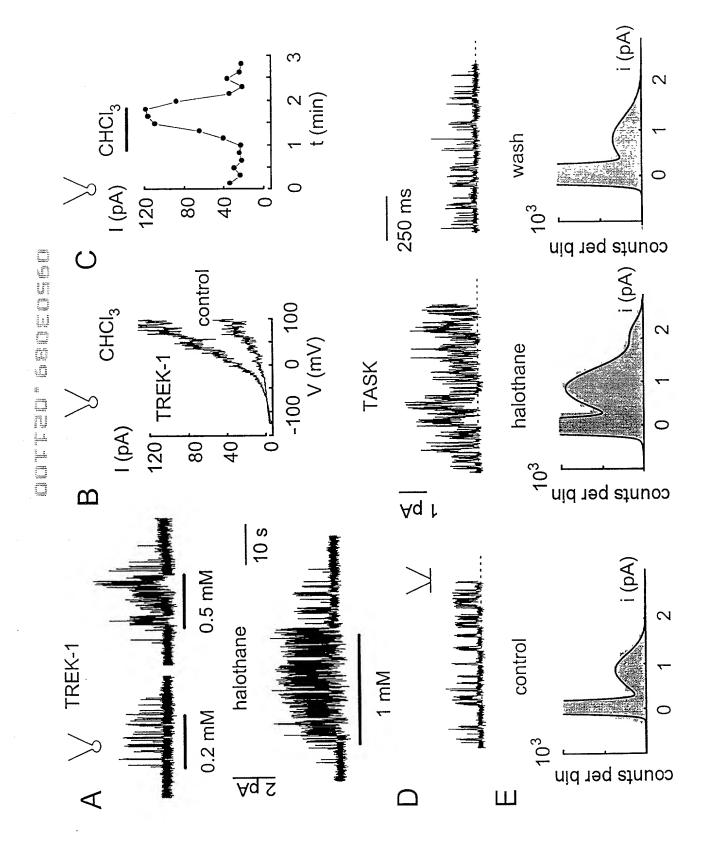


Figure 4A-4E

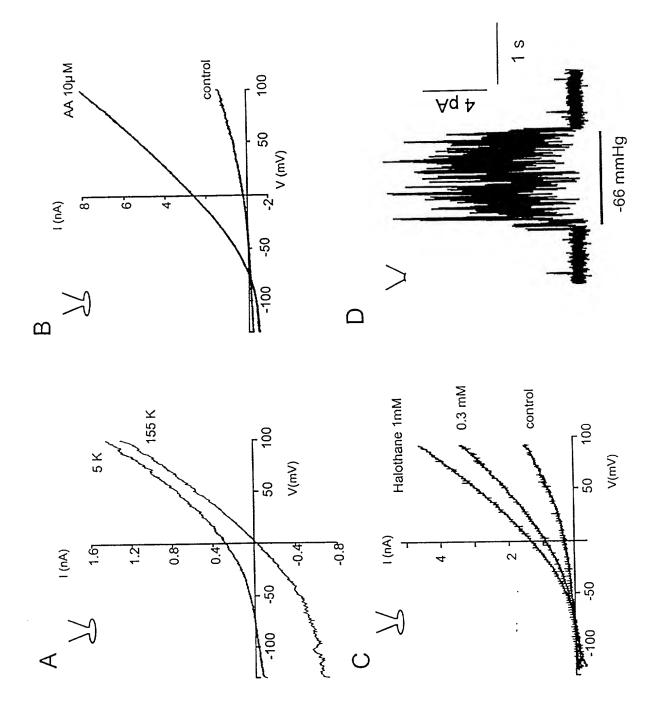


FIGURE 5A-D

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Filing Date	Febr	uary 11, 2000					
Group Art Unit	Not						
Examiner Name	Not :	yet assigned					
	First Named Inventor COMPL Application Number Filing Date Group Art Unit	Application Number Not : Filing Date Febr Group Art Unit Not :	Amanda J. PATEL COMPLETE IF KNOWN Application Number Not yet known Filing Date February 11, 2000 Group Art Unit Not yet assigned				

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My residence, post office ac	dress, and cit	zenship are as stated	below next to my name		
I believe I am the original, fa	st and sole in	ventor (if only one nan	ne is listed below) or an origi	nal, first and joint inve	ntor (if plural
names are listed below) of t	ne subject ma	tter which is claimed a	nd for which a patent is sough	ght on the invention er	htitled:
A METHOD FOR	THE IDEN	NTIFICATION (OF ANESTHETICS		
the specification of which		(Title	of the Invention)		
Is attached hereto		(1.1.2	,		
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was filed on (MM/D	D/YYYY)		as Unit	ed States Application	Number or PCT International
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[Page 1 of 2]

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Amanda J.				PA	TEL	·								
Inventor's Signature								Date						
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ADDITIONAL INVENTOR(S) Supplemental Sheet Page <u>1</u> of <u>2</u>

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Eric				HONORE								
Inventor's Signature							Date					
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Michel				LAZ	DUSKI					
Inventor's Signature		1	ı					Date		
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Glu Asp Thr Phe Ile Lys Trp Asn Val Ser Gln Thr Lys Ile Arg Ile
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                                               205
Ile Ser Thr Ile Ile Phe Ile Leu Phe Gly Cys Val Leu Phe Val Ala
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                                            220
Leu Pro Ala Ile Ile Phe Lys His Ile Glu Gly Trp Ser Ala Leu Asp
                    230
                                       235
Ala Ile Tyr Phe Val Val Ile Thr Leu Thr Thr Ile Gly Phe Gly Asp
                                    250
Tyr Val Ala Gly Gly Ser Asp Ile Glu Tyr Leu Asp Phe Tyr Lys Pro
           260
                                265
Val Val Trp Phe Trp Ile Leu Val Gly Leu Ala Tyr Phe Ala Ala Val
                           280
Leu Ser Met Ile Gly Asp Trp Leu Arg Val Ile Ser Lys Lys Thr Lys
                       295
Glu Glu Val Gly Glu Phe Arg Ala His Ala Ala Glu Trp Thr Ala Asn
                                        315
Val Thr Ala Glu Phe Lys Glu Thr Arg Arg Arg Leu Ser Val Glu Ile
                                    330
Tyr Asp Lys Phe Gln Arg Ala Thr Ser Ile Lys Arg Lys Leu Ser Ala
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                                345
Glu Leu Ala Gly Asn His Asn Gln Glu Leu Thr Pro Cys Arg Arg Thr
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Leu Ser Val Asn His Leu Thr Asn Glu Arg Asp Val Leu Pro Pro Leu
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gga ttt gga aac atc tcc cca cga act gaa ggt gga aaa ata ttc tgc

960

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gly aaa	gtt Val	ggt Gly	gat Asp	cag Gln 180	cta Leu	gga Gly	act Thr	ata Ile	ttt Phe 185	gga Gly	aaa Lys	gga Gly	att Ile	gcc Ala 190	aaa Lys	1056
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gac Asp	tac Tyr	gtg Val	gca Ala	ggt Gly 260	gga Gly	tca Ser	gac Asp	att Ile	gaa Glu 265	tat Tyr	ctg Leu	gac Asp	ttc Phe	tac Tyr 270	гая	1296
cct Pro	gtg Val	gtg Val	tgg Trp 275	ttc Phe	tgg Trp	atc Ile	ctc Leu	gtt Val 280	Gly 999	ctg Leu	gcc Ala	tac Tyr	ttt Phe 285	gca Ala	gct Ala	1344
gtt Val	ctg Leu	agc Ser 290	atg Met	att Ile	gly ggg	gac Asp	tgg Trp 295	Leu	cgg Arg	gtg Val	atc Ile	tct Ser 300	aag Lys	aag Lys	acg Thr	1392
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aat Asn 320	Val	acg Thr	gcc Ala	gag Glu	ttc Phe 325	Lys	gaa Glu	acg Thr	agg Arg	agg Arg 330	Arc	r ctg r Leu	agc Ser	gtg Val	g gag Glu 335	1488
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acc Thr	cto Lev	g tct 1 Sei 370	. Val	g aad L Asi	c cad n His	c ctg Lei	g aco i Thi 375	s Ser	gag Gli	g agg	g gaa g Glu	a gto 1 Val 380	ьег	g cci	t ccc o Pro	1632
tt <u>e</u> Lei	g cto 1 Let 38	ı Lys	g gct s Ala	gaq a Gli	g ago ı Sei	ato r Ile 39	e Ty	t ctg	g aad 1 Asi	c ggt n Gly	cto Y Let 39	ı Tnı	a cca c Pro	a ca o Hi	c tgt s Cys	1680
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Val	Glu	Ser	Asp	Ser	Ala	Ile	Asn	Val	Met	Lys	Trp	Lys	Thr	Val	Ser
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Thr	Ile	Phe	Leu	Val	Val	Val	Leu	Tyr	Leu	Ile	Ile	Gly	Ala	Ala	Val
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Val	Ile	Gln	Lys	Gln	Thr	Phe	Ile	Ala	Gln	His	Ala	Cys	Val	Asn	Ser
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Thr	Glu	Leu	Asp	Glu	Leu	Ile	Gln	Gln	Ile	Val	Ala	Ala	Ile	Asn	Ala
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Leu	Gly	Ser	Ser	Phe	Phe	Phe	Ala	Gly	Thr	Val	Ile	Thr	Thr	Ile	Gly
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Phe	Gly	Asn	Ile	Ser	Pro	Arg	Thr	Glu	Gly	Gly	Lys	Ile	Phe	Cys	Ile
145					150					155					160
Ile	Tyr	Ala	Leu	Leu	Gly	Ile	Pro	Leu	Phe	Gly	Phe	Leu	Leu	Ala	Gly
				165					170					175	
Val	Gly	Asp	Gln	Leu	Gly	Thr	Ile	Phe	Gly	Lys	Gly	Ile	Ala	Lys	Val
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Glu	Asp	Thr	Phe	e Ile	Lys	Trp	Asn	Val	Ser	Gln	Thr	Lys	Ile	Arg	Ile
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Ile Ser Thr Ile Ile Phe Ile Leu Phe Gly Cys Val Leu Phe Val Ala

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Ala	Ile	Tyr	Phe	Val	Val	Ile	Thr	Leu	Thr	Thr	Ile	Gly	Phe	Gly	Asp
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Tyr	Val	Ala	Gly	Gly	Ser	Asp	Ile	Glu	Tyr	Leu	Asp	Phe	Tyr	Lys	Pro
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Val	Val	Trp	Phe	Trp	Ile	Leu	Val	Gly	Leu	Ala	Tyr	Phe	Ala	Ala	Val
		275					280					285			
Leu	Ser	Met	Ile	Gly	Asp	Trp	Leu	Arg	Val	Ile	Ser	Lys	Lys	Thr	Lys
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Val	Thr	Ala	Glu	Phe	Lys	Glu	Thr	Arg	Arg	Arg	Leu	Ser	Val	Glu	Ile
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Val	Val	Leu	Arg	Leu	Lys	Pro	His	Lys	Ala	Gly	Val	Gln	Trp	Arg	Phe
65					70					75					80
Ala	Gly	Ser	Phe	Tyr	Phe	Ala	Ile	Thr	Val	Ile	Thr	Thr	Ile	Gly	Tyr
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Gly	His	Ala	Ala	Pro	Ser	Thr	Asp	Gly	Gly	Lys	Val	Phe	Cys	Met	Phe
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Ala	Phe	Ser	His	Tyr	Glu	His	Trp	Thr	Phe	Phe	Gln	Ala	Tyr	Tyr	Tyr
			180					185					190		
Cys	Phe	Ile	Thr	Leu	Thr	Thr	Ile	Gly	Phe	Gly	Asp	Tyr	Val	Ala	Leu
		195					200					205			
Gln	Lys	Asp	Gln	Ala	Leu	Gln	Thr	Gln	Pro	Gln	Tyr	Val	Ala	Phe	Ser
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Phe	Val	Tyr	Ile	Leu	Thr	Gly	Leu	Thr	Val	Ile	Gly	Ala	Phe	Leu	Asn
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Asp	Ala	Glu	His	Arg	Ala	Leu	Leu	Thr	Arg	Asn	Gly	Gln	Ala	Gly	Gly
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Gly	Gly	Gly	Gly	Gly	Ser	Ala	His	Thr	Thr	Asp	Thr	Ala	Ser	Ser	Thr
		275					280					285			
Ala	Ala	Ala	Gly	Gly	Gly	Gly	Phe	Arg	Asn	Val	Tyr	Ala	Glu	Val	Leu
	290					295					300				
His	Phe	Gln	Ser	Met	Cys	Ser	Cys	Leu	Trp	Tyr	Lys	Ser	Arg	Glu	Lys
305					310					315					320
Leu	Gln	Tyr	Ser	Ile	Pro	Met	Ile	Ile	Pro	Arg	Asp	Leu	Ser	Thr	Ser
				325					330					335	
Asp	Thr	Cys	Val	Glu	Gln	Ser	His	Ser	Ser	Pro	Gly	Gly	Gly	Gly	Arg
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Tyr	Ser	Asp	Thr	Pro	Ser	Arg	Arg	Cys	Leu	Cys	Ser	Gly	Ala	Pro	Arg
		355					360					365			
Ser	Ala	Ile	Ser	Ser	Val	Ser	Thr	Gly	Leu	His	Ser	Leu	Ser	Thr	Phe
	370					375					380				
Arg	Gly	Leu	Met	Lys	Arg	Arg	Ser	Ser	Val						
385					390										